		Serla	CRF Errors Corrected by the STIC Systems Branch / 6/1/  Changed a file from non-ASCII to ASCII  Changed a file from non-ASCII to ASCII  Changed a file from non-ASCII to ASCII
•			, ————————————————————————————————————
PEC 0 7 2001  DEC 0 7 2001			Changed the margins in cases where the sequence text was "wrapped" down to the next line
	<b>/</b>		Edited a format error in the Current Application Data section, specifically:
	<b>1</b>		Edited the Current Application Data section with the actual current number. The number input applicant was the prior application data; or other
			Added the mandatory heading and subheadings for "Current Application Data".
	•		Edited the "Number of Sequences" field. The applicant spelled out a number instead of using
			Changed the spelling of a mandatory field (the headings or subheadings), specifically;
	•		Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited
			Inserted or corrected a nucleic number at the end of a nucleic line. SEO D NO: edited:
	٥		Corrected subheading placement. All responses must be on the same line as each subheading applicant placed a response below the subheading, this was moved to its appropriate place.
	$\Omega$		Inserted colons after headings/subheadings. Headings edited included:
	EVE		Deleted extra, invalid, headings used by an applicant, specifically:
	Ü		Deleted: non-ASCII *garbage* at the beginning/end of files; secretary initials/filename a page numbers throughout text; other invalid text, such as
			Inserted mandatory headings, specifically: Header Geld 220 requeste
			Corrected an obvious error in the response, specifically:
			Edited identifiers where upper case is used but lower case is required, or vice versa.
	. 1		Corrected an error in the Number of Sequences field, specifically:
		•	·
	. [		A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
			Deleted ending stop codon in amino acid sequences and adjusted the *(A)Length: field according due to a Patentin bug). Sequences corrected:
	[		Other:
		•	
		-	<u> </u>

\*Examiner: The above corrections must be communicated to the applicant in the first ( Action. DO NOT send a copy of this form.

RAW SEQUENCE LISTING

4 <110> APPLICANT: Anderson, George P.

PATENT APPLICATION: US/09/811,824

DATE: 11/13/2001 TIME: 10:32:52

Input Set : A:\SEQUENCE LISTING.pto.mph.txt
Output Set: N:\CRF3\11092001\1811824.raw

```
Mattoussi, Hedi
 6
         Mauro, J. Matthew
 7
         Bawendi, Moungi G.
         Sundar, Vikram C.
10 <120> TITLE OF INVENTION: INORGANIC PARTICLE CONJUGATES
12 <130> FILE REFERENCE: 01997-282001
14 <140> CURRENT APPLICATION NUMBER: US 09/811,824
15 <141> CURRENT FILING DATE: 2001-03-20
                                                                 ENTEREC
17 <150> PRIOR APPLICATION NUMBER: US 60/190,766
18 <151> PRIOR FILING DATE: 2000-03-20
20 <160> NUMBER OF SEQ ID NOS: 7
22 <170> SOFTWARE: FastSEQ for Windows Version 4.0
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 21
26 <212> TYPE: DNA
27 <213> ORGANISM: Artificial Sequence
29 <220> FEATURE:
30 <223> OTHER INFORMATION: Synthetically generated primer
32 <400> SEQUENCE: 1
33 tgcggtggct cagctcagtt g
                                                                           21
35 <210> SEQ ID NO: 2
36 <211> LENGTH: 33
37 <212> TYPE: DNA
38 <213> ORGANISM: Artificial Sequence
40 <220> FEATURE:
41 <223> OTHER INFORMATION: Synthetically generated primer
43 <400> SEQUENCE: 2
44 gctctagatt aatccccacc ctgggcgagt ttc
                                                                           33
46 <210> SEQ ID NO: 3
47 <211> LENGTH: 33
48 <212> TYPE: DNA
49 <213> ORGANISM: Artificial Sequence
51 <220> FEATURE:
52 <223> OTHER INFORMATION: Synthetically generated primer
54 <400> SEQUENCE: 3
55 gctctagatg aatccccacc ctgggcgagt ttc
                                                                           33
58 <210> SEO ID NO: 4
59 <211> LENGTH: 34
60 <212> TYPE: DNA
61 <213> ORGANISM: Artificial Sequence
63 <220> FEATURE:
64 <223> OTHER INFORMATION: Synthetically generated primer,
66 <400> SEQUENCE: 4
67 ctagcggtca ccaccaccac caccactgac tgca
69 <210> SEO ID NO: 5
70 <211> LENGTH: 26
```

NuEnorel

RAW SEQUENCE LISTING DATE: 11/13/2001 PATENT APPLICATION: US/09/811,824 TIME: 10:32:52

Input Set : A:\SEQUENCE LISTING.pto.mph.txt
Output Set: N:\CRF3\11092001\I811824.raw

```
71 <212> TYPE: DNA
72 <213> ORGANISM: Artificial Sequence
74 <220> FEATURE:
75 <223> OTHER INFORMATION: Synthetically generated primer -
77 <400> SEQUENCE: 5
78 gtcagtggtg gtggtggtgg tgaccg
                                                                            26
80 <210> SEQ ID NO: 6
81 <211> LENGTH: 105
82 <212> TYPE: DNA
83 <213> ORGANISM: Artificial Sequence,
85 <220> FEATURE:
86 <223> OTHER INFORMATION: Synthetically generated primer
88 <400> SEQUENCE: 6
89 tcagctcagt tgaaaaaaaa attgcaagca ctgaagaaaa agaacgctca gctgaagtgg
90 aaacttcaag ccgtcaagaa gaaactcgcc cagggtgggg attca
                                                                          105
92 <210> SEQ ID NO: 7
93 <211> LENGTH: 35
94 <212> TYPE: PRT
95 <213> ORGANISM: Artificial Sequence
97 <220> FEATURE:
98 <223> OTHER INFORMATION: Synthetically generated peptide.
100 <400> SEQUENCE: 7
101 Ser Ala Gln Leu Lys Lys Lys Leu Gln Ala Leu Lys Lys Lys Asn Ala
102
                                          10
103 Gln Leu Lys Trp Lys Leu Gln Ala Leu Lys Lys Lys Leu Ala Gln Gly
                                     25
105 Gly Asp Ser
106
             35
```

VERIFICATION SUMMARY

VERIFICATION SUMMARYDATE: 11/13/2001PATENT APPLICATION: US/09/811,824TIME: 10:32:53

Input Set : A:\SEQUENCE LISTING.pto.mph.txt Output Set: N:\CRF3\11092001\1811824.raw

OIPE

RAW SEQUENCE LISTING

DATE: 11/13/2001

PATENT APPLICATION: US/09/811,824

TIME: 10:25:10

Input Set : A:\SEQUENCE LISTING.pto.mph.txt Output Set: N:\CRF3\11092001\I811824.raw

- 4 <110> APPLICANT: Anderson, George P.
- Mattoussi, Hedi
- Mauro, J. Matthew 6
- Bawendi, Moungi G.
- Sundar, Vikram C.
- 10 <120> TITLE OF INVENTION: INORGANIC PARTICLE CONJUGATES
- 12 <130> FILE REFERENCE: 01997-282001
- 14 <140> CURRENT APPLICATION NUMBER: US 09/811,824
- 15 <141> CURRENT FILING DATE: 2001-03-20
- 17 <150> PRIOR APPLICATION NUMBER: US 60/190,766
- 18 <151> PRIOR FILING DATE: 2000-03-20
- 20 <160> NUMBER OF SEQ ID NOS: 7
- 22 <170> SOFTWARE: FastSEQ for Windows Version 4.0

## ERRORED SEQUENCES

46 <210> SEQ ID NO: 3

47 <211> LENGTH: 33

48 <212> TYPE: DNA

49 <213> ORGANISM: Artificial Sequence

E--> 50 (200)> W--> 51 (220) FEATURE:

51 COTHER INFORMATION: Synthetically generated primer

53 <400> SEQUENCE: 3

54 gctctagatg aatccccacc ctgggcgagt ttc

33

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/811,824

DATE: 11/13/2001 TIME: 10:25:11

Input Set : A:\SEQUENCE LISTING.pto.mph.txt
Output Set: N:\CRF3\11092001\1811824.raw

L:50 M:250 E: Invalid Numeric Identifier, INVALID IDENTIFIER

L:51 M:258 W: Mandatory Feature missing, <220> FEATURE:

STATISTICS SUMMARY

PATENT APPLICATION: US/09/811,824

DATE: 11/13/2001 TIME: 10:25:11

Input Set : A:\SEQUENCE LISTING.pto.mph.txt
Output Set: N:\CRF3\11092001\1811824.raw

Application Serial Number: US/09/811,824

Alpha or Numeric: Numeric

Application Class:

Application File Date: 03-20-2001

Art Unit: OIPE

Software Application: FastSeq Total Number of Sequences: 7

Total Nucleotides: 252
Total Amino Acids: 35
Number of Errors: 1
Number of Warnings: 1
Number of Corrections: 0

## MESSAGE SUMMARY

250 E: 1 (Invalid Numeric Identifier)
258 W: 1 (Mandatory Feature missing)

OIPE

RAW SEQUENCE LISTING

DATE: 11/06/2001 TIME: 10:57:40

PATENT APPLICATION: US/09/811,824

Input Set : A:\SEQUENCE LISTING.TXT Output Set: N:\CRF3\11062001\1811824.raw

- 4 <110> APPLICANT: Anderson, George P.
- Mattoussi, Hedi
- Mauro, J. Matthew
- Bawendi, Moungi G.
- Sundar, Vikram C.
- 10 <120> TITLE OF INVENTION: INORGANIC PARTICLE CONJUGATES
- 12 <130> FILE REFERENCE: 01997-282001
- 14 <140> CURRENT APPLICATION NUMBER: US 09/811,824
- 15 <141> CURRENT FILING DATE: 2001-03-20
- 17 <150> PRIOR APPLICATION NUMBER: US 60/190,766
- 18 <151> PRIOR FILING DATE: 2000-03-20
- 20 <160> NUMBER OF SEQ ID NOS: 7
- 22 <170> SOFTWARE: FastSEQ for Windows Version 4.0

## **ERRORED SEQUENCES**

- 46 <210> SEQ ID NO: 3
- 47 <211> LENGTH: 33
- 48 <212> TYPE: DNA
- 49 <213> ORGANISM: Artificial Sequence La leve

E > 51 (200) 52 <220> FEATURE:

52 <223> OTHER INFORMATION: Synthetically generated primer

54 <400> SEQUENCE: 3

55 gctctagatg aatccccacc ctgggcgagt ttc

33

Field 220 to mandatury pertur to field 221, 222 or 223

VERIFICATION SUMMARY

DATE: 11/06/2001 TIME: 10:57:41

PATENT APPLICATION: US/09/811,824

Input Set : A:\SEQUENCE LISTING.TXT Output Set: N:\CRF3\11062001\1811824.raw

L:51 M:250 E: Invalid Numeric Identifier, INVALID IDENTIFIER

L:52 M:258 W: Mandatory Feature missing, <220> FEATURE:

STATISTICS SUMMARY

DATE: 11/06/2001

PATENT APPLICATION: US/09/811,824

TIME: 10:57:41

Input Set : A:\SEQUENCE LISTING.TXT Output Set: N:\CRF3\11062001\1811824.raw

Application Serial Number: US/09/811,824

Alpha or Numeric: Numeric

Application Class:

Application File Date: 03-20-2001

Art Unit: OIPE

Software Application: FastSeq Total Number of Sequences: 7

Total Nucleotides: 252 Total Amino Acids: 35 Number of Errors: 1 Number of Warnings: 1 Number of Corrections: 0

## MESSAGE SUMMARY

250 E: 1 (Invalid Numeric Identifier) 258 W: 1 (Mandatory Feature missing)